SEQUENCE LISTING

<110> Krolewski, Andrzej S.
      Pezzolesi, Marcus G.
      Nagase, Terumasa

<120> TYPE 2 DIABETES MELLITUS GENES

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<141> 2003-10-28

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<213> Homo sapiens

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Met Pro Ala Lys Ser Ser Lys Met Tyr Gly Thr Leu Arg Lys Gly Ser
    50                  55                  60
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Val Cys Ala Asp Pro Lys Pro Gln Gln Val Lys Lys Ile Phe Glu Ala
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Leu Lys Arg Gly Leu Lys Glu Tyr Leu Cys Val Gln Gln Ala Glu Leu
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Asp His Leu Ser Gly Arg His Lys Asp Thr Arg Arg Asn Ser Arg Leu
            100                 105                 110
Ala Phe Tyr Tyr Asp Leu Asp Lys Gln Thr Arg Cys Val Glu Arg His
        115                 120                 125
Ile Arg Lys Met Glu Phe His Ile Ser Lys Val Asp Glu Leu Tyr Glu
    130                 135                 140
Asp Tyr Cys Ile Gln Cys Arg Leu Arg Asp Gly Ala Ser Ser Met Gln
145                 150                 155                 160
Arg Ala Phe Ala Arg Cys Pro Pro Ser Arg Ala Ala Arg Glu Ser Leu
                165                 170                 175
Gln Glu Leu Gly Arg Ser Leu His Glu Cys Ala Glu Asp Met Trp Leu
            180                 185                 190
Ile Glu Gly Ala Leu Glu Val His Leu Gly Glu Phe His Ile Arg Met
            195                 200                 205
Lys Gly Leu Val Gly Tyr Ala Arg Leu Cys Pro Gly Asp His Tyr Glu
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Val Leu Met Arg Leu Gly Arg Gln Arg Trp Lys Leu Lys Gly Arg Ile
225                 230                 235                 240
Glu Ser Asp Asp Ser Gln Thr Trp Asp Glu Glu Glu Lys Ala Phe Ile
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Pro Thr Leu His Glu Asn Leu Asp Ile Lys Val Thr Glu Leu Arg Gly
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Leu Gly Ser Leu Ala Val Gly Ala Val Thr Cys Asp Ile Ala Asp Phe
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Phe Thr Thr Arg Pro Gln Val Ile Val Val Asp Ile Thr Glu Leu Gly
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Thr Ile Lys Leu Gln Leu Glu Val Gln Trp Asn Pro Phe Asp Thr Glu
305                 310                 315                 320
Ser Phe Leu Val Ser Pro Ser Pro Thr Gly Lys Phe Ser Met Gly Ser
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Arg Lys Gly Ser Leu Tyr Asn Trp Thr Pro Pro Ser Thr Pro Ser Phe
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            355                 360                 365
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Gly Pro His Ala Ser Ile Glu Glu Glu Ala Arg Glu Asp Pro Leu Pro
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Pro Gly Leu Leu Pro Glu Met Ala His Leu Ser Gly Gly Pro Phe Ala
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Glu Gln Pro Gly Trp Arg Asn Leu Gly Gly Glu Ser Pro Ser Leu Pro
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Gln Gly Ser Leu Phe His Ser Gly Thr Ala Ser Ser Ser Gln Asn Gly
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His Glu Glu Gly Ala Thr Gly Asp Arg Glu Asp Gly Pro Gly Val Ala
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Leu Glu Gly Pro Leu Gln Glu Val Leu Glu Leu Leu Arg Pro Thr Asp
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Asp Phe Ala Leu Asp Glu Leu Ser Leu Phe Gly Gly Ser Gln Gly Leu
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Arg Lys Asp Arg Pro Leu Pro Pro Ser Ser Leu Lys Ala Ser Ser
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Arg Glu Leu Thr Ala Gly Ala Pro Glu Leu Asp Val Leu Leu Met Val
        610                     615                     620
His Leu Gln Val Cys Lys Ala Leu Leu Gln Lys Leu Ala Ser Pro Asn
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Leu Ser Arg Leu Val Gln Glu Cys Leu Leu Glu Glu Val Ala Gln Gln
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Lys His Val Leu Glu Thr Leu Ser Val Leu Asp Phe Glu Lys Val Gly
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Lys Ala Thr Ser Ile Glu Glu Ile Ile Pro Gln Ala Ser Arg Thr Lys
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Gly Cys Leu Lys Leu Trp Arg Gly Cys Thr Gly Pro Gly Arg Val Leu
        690                     695                     700
Ser Cys Pro Ala Thr Thr Leu Leu Asn Gln Leu Lys Lys Thr Phe Gln
705                     710                  .  715                     720
His Arg Val Arg Gly Lys Tyr Pro Gly Gln Leu Glu Ile Ala Cys Arg
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Arg Leu Leu Glu Gln Val Val Ser Cys Gly Gly Leu Leu Pro Gly Ala
            740                     745                     750
Gly Leu Pro Glu Glu Gln Ile Ile Thr Trp Phe Gln Phe His Ser Tyr
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Leu Gln Arg Gln Ser Val Ser Asp Leu Glu Lys His Phe Thr Gln Leu
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Thr Lys Glu Val Thr Leu Ile Glu Glu Leu His Cys Ala Gly Gln Ala
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Lys Val Val Arg Lys Leu Gln Gly Lys Arg Leu Gly Gln Leu Gln Pro
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Leu Pro Gln Thr Leu Arg Ala Trp Ala Leu Leu Gln Leu Asp Gly Thr
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Pro Arg Val Cys Arg Ala Ala Ser Ala Arg Leu Ala Gly Ala Val Arg
            835                     840                     845
Asn Arg Ser Phe Arg Glu Lys Ala Leu Leu Phe Tyr Thr Asn Ala Leu
    850                     855                     860
Ala Glu Asn Asp Ala Arg Leu Gln Gln Ala Ala Cys Leu Ala Leu Lys
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His Leu Lys Gly Ile Glu Ser Ile Asp Gln Thr Ala Ser Leu Cys Gln
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Ser Asp Leu Glu Ala Val Arg Ala Ala Ala Arg Glu Thr Thr Leu Ser
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Phe Gly Glu Lys Gly Arg Leu Ala Phe Glu Lys Met Asp Lys Leu Cys
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Ile Phe
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<211> 2278
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<213> Homo sapiens

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atgggagcag acgcttgctg gcggcaatag ggaaagtgag gcagctgcaa ggagggcggc     300
gggactgcac tcgagtgtcc agacctgctc gatggtgacc accatgtcgg tgaggttgcg     360
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<212> PRT
<213> Homo sapiens

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Met Pro Ala Lys Ser Ser Lys Met Tyr Gly Thr Leu Arg Lys Gly Ser
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Val Cys Ala Asp Pro Lys Pro Gln Gln Val Lys Lys Ile Phe Glu Ala
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Asp His Leu Ser Gly Arg His Lys Asp Thr Arg Arg Asn Ser Arg Leu
                 100                   105                   110
Ala Phe Tyr Tyr Asp Leu Asp Lys Gln Thr Arg Cys Val Glu Arg His
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Ile Arg Lys Met Glu Phe His Ile Ser Lys Val Asp Glu Leu Tyr Glu
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Asp Tyr Cys Ile Gln Cys Arg Leu Arg Asp Gly Ala Ser Ser Met Gln
145                   150                   155                   160
Arg Ala Phe Ala Arg Cys Pro Pro Ser Arg Ala Ala Arg Glu Ser Leu
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Gln Glu Leu Gly Arg Ser Leu His Glu Cys Ala Glu Asp Met Trp Leu
             180                   185                   190
Ile Glu Gly Ala Leu Glu Val His Leu Gly Glu Phe His Ile Arg Met
             195                   200                   205
Lys Gly Leu Val Gly Tyr Ala Arg Leu Cys Pro Gly Asp His Tyr Glu
         210                   215                   220
Val Leu Met Arg Leu Gly Arg Gln Arg Trp Lys Leu Lys Gly Arg Ile
225                   230                   235                   240
Glu Ser Asp Asp Ser Gln Thr Trp Asp Glu Glu Glu Lys Ala Phe Ile
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Pro Thr Leu His Glu Asn Leu Asp Ile Lys Val Thr Glu Leu Arg Gly
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Leu Gly Ser Leu Ala Val Gly Ala Val Thr Cys Asp Ile Ala Asp Phe
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Phe Thr Thr Arg Pro Gln Val Ile Val Val Asp Ile Thr Glu Leu Gly
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Arg Lys Gly Ser Leu Tyr Asn Trp Thr Pro Pro Ser Thr Pro Ser Phe
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Arg Glu Arg Tyr Tyr Leu Ser Val Leu Gln Gln Pro Thr Gln Gln Ala
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Leu Leu Leu Gly Gly Pro Arg Ala Thr Ser Ile Leu Ser Tyr Leu Ser
  · 370                   375                   380
Asp Ser Asp Leu Arg Gly Pro Ser Leu Arg Ser Gln Ser Gln Glu Leu
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Pro Glu Met Asp Ser Phe Ser Ser Glu Asp Pro Arg Asp Thr Glu Thr
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Ser Thr Ser Ala Ser Thr Ser Asp Val Gly Phe Leu Pro Leu Thr Phe
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Gly Pro His Ala Ser Ile Glu Glu Glu Ala Arg Glu Asp Pro Leu Pro
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Gln Gly Ser Leu Phe His Ser Gly Thr Ala Ser Ser Ser Gln Asn Gly
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His Glu Glu Gly Ala Thr Gly Asp Arg Glu Asp Gly Pro Gly Val Ala
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545   550   555   560
Phe Ala Asp Gly Met Met Thr Gly Ser Arg Gly Leu Trp Gly His Ala
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Ala Trp Ala Asp Ile Leu Ala Ser Pro Leu Arg Asp Leu Gly Gly Pro
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Cys Leu Ser Gly Pro Trp Phe Pro His Leu Ala Ser Gly Asp Asn Asn
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<212> DNA
<213> Homo sapiens

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<210> 9
<211> 403
<212> DNA
<213> Homo sapiens

<400> 9
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<210> 10
<211> 399
<212> DNA
<213> Homo sapiens

<400> 10
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<210> 11
<211> 401
<212> DNA
<213> Homo sapiens

<400> 11
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ctccatgttt  ttttaaagct  ttatgcacaa  acatgatcat  aagacatgat  ttatgataaa         300
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<210> 12
<211> 401
<212> DNA
<213> Homo sapiens

<400> 12
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agggaaagcg  ggcatcccct  cacatccatg  ggatctatgt  gtgggttgta  tcaagagtct         180
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ctccatgttt  ttttaaagct  ttatgcacaa  acatgatcat  aagacatgat  ttatgataaa         300
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<210> 13
<211> 401
<212> DNA
<213> Homo sapiens

<400> 13
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cctagagtca  cacagctgac  tgaagagtgt  gctgcaactc  caggacttgt  ctcccttacc         300
tccccacaaa  gagtgtgtat  ctctgagccc  agcccagcca  cagcctccac  tctgggcccc         360
gattaactct  ggctattagg  aaggcagaag  aggctccccg  a                              401
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<210> 14
<211> 401
<212> DNA
<213> Homo sapiens

<400> 14
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aaggaacctg  gccttctgag  gggaggtccc  acggggcaga  ggcacagctg  ggatcacagc         120
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cctagagtca cacagctgac tgaagagtgt gctgcaactc caggacttgt ctcccttacc    300
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gattaactct ggctattagg aaggcagaag aggctccccg a                        401
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<210> 15
<211> 401
<212> DNA
<213> Homo sapiens

<400> 15
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tgcactccat caggctctcg gccgaggtgt gctcctgccg tgctctgcag ggctgtggac    300
gaagtggcca gacctgaggg caacaccggg ccccacccac ccgactggga cactggccag    360
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<210> 16
<211> 401
<212> DNA
<213> Homo sapiens

<400> 16
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gtggtggcca ctgggtcact gctggaagga ggtgcactca ccggagaccc tgggagcccc    180
caaacaggga cagctcatcc ggggcgaagt cggcattgag gaaggcgaag ctctccagga    240
tgcactccat caggctctcg gccgaggtgt gctcctgccg tgctctgcag ggctgtggac    300
gaagtggcca gacctgaggg caacaccggg ccccacccac ccgactggga cactggccag    360
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<210> 17
<211> 401
<212> DNA
<213> Homo sapiens

<400> 17
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cccctcgatg agccacatgt cctgcaaagc cccggaggtg gctcagctgg ctgcctgggg    180
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<210> 18
<211> 401
<212> DNA
<213> Homo sapiens

<400> 18
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cccctcgatg agccacatgt cctgcaaagc cccggaggtg gctcagctgg ctgcctgggg    180
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gcctccttgg ggcccagaac acctcctcca gcccccactg gcccagctct cgatgtcccc     300
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ggcgcactcg tgcaggctgc ggcccagctc ctgcaggctc t                         401

<210> 19
<211> 416
<212> DNA
<213> Homo sapiens

<400> 19
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tttgcaaagt tgcctgggcc acccacctgc cccgcttgcc cctgccaccc tcctacaggt     180
cctaactcag agaatggggc ttagtgccgg gccggcccct caccatccct gaggaaggct     240
catcgcagag actcagcctt cccattccta aaatggggag gagacccagg ttttctgccc     300
atcaggcagc caggaagatg caatgaggca cagtcattct catccagcca ggcccagccc     360
acctcactca ccgtatgcag actcaccttg tccaggtcat aatagaaagc ctgtga         416

<210> 20
<211> 400
<212> DNA
<213> Homo sapiens

<400> 20
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tttgcaaagt tgcctgggcc acccacctgc cccgcttgcc cctgccaccc tcctacaggt     180
cctaactcag agaatggggc ccctcaccat ccctgaggaa ggctcatcgc agagactcag     240
ccttcccatt cctaaaatgg ggaggagacc caggtttct gcccatcagg cagccaggaa      300
gatgcaatga ggcacagtca ttctcatcca gccaggccca gcccacctca ctcaccgtat     360
gcagactcac cttgtccagg tcataataga aagcctgtga                           400

<210> 21
<211> 401
<212> DNA
<213> Homo sapiens

<400> 21
cagcggcaga ggccactgtg acatacccaa gatgtgacac ctgacccact ttcctggcat     60
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taaaagaaat gcacctccat acactgccca ccttagcatt acttctagaa ccgagagaca     180
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gcaacgtgaa gaggctattc cataaacctc tagttctgag aaagagtcac accgtgacat     300
aggctagaag gaacgcaggg ttcatctttt actcctggcc aaggctatct gggtgggaag     360
caggcaggga ggggtctcac cagcctggaa ttcctcctgg t                         401

<210> 22
<211> 401
<212> DNA
<213> Homo sapiens

<400> 22
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taaaagaaat gcacctccat acactgccca ccttagcatt acttctagaa ccgagagaca     180
gtgtgacatg ggcctaaaac gtgtgaactg ctgtacgtgc caaagtgaag ttaactcagt     240
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gcaacgtgaa  gaggctattc  cataaacctc  tagttctgag  aaagagtcac  accgtgacat      300
aggctagaag  gaacgcaggg  ttcatctttt  actcctggcc  aaggctatct  gggtgggaag      360
caggcaggga  ggggtctcac  cagcctggaa  ttcctcctgg  t                           401
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<210> 23
<211> 401
<212> DNA
<213> Homo sapiens

<400> 23
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ctgcacagac  cgaccccttc  cgcagcgtgc  cgtacatctt  ggaggatttt  gcaggcattc      360
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<210> 24
<211> 401
<212> DNA
<213> Homo sapiens

<400> 24
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ctgcacagac  cgaccccttc  cgcagcgtgc  cgtacatctt  ggaggatttt  gcaggcattc      360
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<210> 25
<211> 401
<212> DNA
<213> Homo sapiens

<400> 25
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cttaagccag  ctggagttgg  gagtgtggac  tggatgatcc  taaaaactgc  ctttcagtgg      180
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tacgaggcca  gggcaccctc  atcacagcac  agagctggtt  tccctggcat  ctaagcctct      300
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<210> 26
<211> 401
<212> DNA
<213> Homo sapiens

<400> 26
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<210>  27
<211>  401
<212>  DNA
<213>  Homo sapiens


<400>  27
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tgccagccgc  acacctgctt  tctgtccctt  aaagctcatt  cccacccagg  acatctgcac      180
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tccacttctt  ctgtcccttc  ctgcctccac  tccccatgcc  cctgtctcgt  caggctctcc      300
caggagacca  tgggtgccct  cccccacccc  cagttcagtt  ccctcacagc  actgccacca      360
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<210>  28
<211>  401
<212>  DNA
<213>  Homo sapiens


<400>  28
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tgccagccgc  acacctgctt  tctgtccctt  aaagctcatt  cccacccagg  acatctgcac      180
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caggagacca  tgggtgccct  cccccacccc  cagttcagtt  ccctcacagc  actgccacca      360
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<210>  29
<211>  401
<212>  DNA
<213>  Homo sapiens


<400>  29
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gagaaggaca  attgctaagc  agttcctccc  gatgcaaagc  tcaaaacaag  ccccaggtcc      360
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<210>  30
<211>  401
<212>  DNA
<213>  Homo sapiens


<400>  30
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gagaataagg  acggagtagg  gagggattcc  caaaggtgag  tagttgagac  gtactccgga      180
gccagcctgg  gcactggagc  tggaaggggc  ttccccggcc  cctccctctg  caccttccca      240
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gagaaggaca  attgctaagc  agttcctccc  gatgcaaagc  tcaaaacaag  ccccaggtcc      360
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<210> 31
<211> 401
<212> DNA
<213> Homo sapiens

<400> 31
```
ccaaggtgtg gctggaggaa gcagagtcta ctcccgctaa gtctgtccgc tcactgctgg    60
ccaaagctgc cctgcgtctc ctccccaccg ccagccagag ggaacctgca atttcacctc   120
atttagaggt aaaacatcta aatttaacgt tatgggcttt tggggctggg tggctttat    180
gcctgagtcc ctcacttagg gctcctttt atccactcaa atgccagcta gggcttagtt    240
tgtttatagg agtttccaaa atagctcctt tggtttcgca tgaaaggaaa tggcaaaata   300
gcccaggaag aggaatgtga gtttacacag aagacagaca ggcgcccgag gaggcttctc   360
tgggaaccag ttcgcctgta ccagagggg cccgagaaag t                        401
```

<210> 32
<211> 401
<212> DNA
<213> Homo sapiens

<400> 32
```
ccaaggtgtg gctggaggaa gcagagtcta ctcccgctaa gtctgtccgc tcactgctgg    60
ccaaagctgc cctgcgtctc ctccccaccg ccagccagag ggaacctgca atttcacctc   120
atttagaggt aaaacatcta aatttaacgt tatgggcttt tggggctggg tggctttat    180
gcctgagtcc ctcacttagg actcctttt atccactcaa atgccagcta gggcttagtt    240
tgtttatagg agtttccaaa atagctcctt tggtttcgca tgaaaggaaa tggcaaaata   300
gcccaggaag aggaatgtga gtttacacag aagacagaca ggcgcccgag gaggcttctc   360
tgggaaccag ttcgcctgta ccagagggg cccgagaaag t                        401
```

<210> 33
<211> 401
<212> DNA
<213> Homo sapiens

<400> 33
```
tacgttagaa ggaccctacg ttagaagggt gaggcgctag ggccatagcc taagggcact    60
gggaaccctg tgggcatgcg cagttcaagc ccatccccgc tccctccagc tgctgtccat   120
ccctgccaca cctgaccatt tgcctaacct agatccttcc tgtcttgcat ttcctcaagc   180
atccggagcc caggactgct gagtcaaccc tctggaatgc ccacaactcc ccacaggcca   240
gccggccttg ggactcccgc acagccacgt gagccggtgg agccgggtct gtttgctagt   300
ggaggctgtt aacagcacgg gaagtggtca agggttcaac aagagatgag ccatctggtc   360
ctccagaggt aaacaattta caagagacac atcaagccgg c                        401
```

<210> 34
<211> 401
<212> DNA
<213> Homo sapiens

<400> 34
```
tacgttagaa ggaccctacg ttagaagggt gaggcgctag ggccatagcc taagggcact    60
gggaaccctg tgggcatgcg cagttcaagc ccatccccgc tccctccagc tgctgtccat   120
ccctgccaca cctgaccatt tgcctaacct agatccttcc tgtcttgcat ttcctcaagc   180
atccggagcc caggactgct cagtcaaccc tctggaatgc ccacaactcc ccacaggcca   240
gccggccttg ggactcccgc acagccacgt gagccggtgg agccgggtct gtttgctagt   300
ggaggctgtt aacagcacgg gaagtggtca agggttcaac aagagatgag ccatctggtc   360
ctccagaggt aaacaattta caagagacac atcaagccgg c                        401
```

<210> 35
<211> 401
<212> DNA
<213> Homo sapiens

<400> 35

```
gggtttcccc caagcccctt tccccctttg cgcctcccac ttctcctaga ttgagagtca      60
gcttggttct ttcctttaca tccattagtg agggtcaggc tcttttgtta tgtttttttt     120
tcttttgtat aacttaatta tttcagggtt cggggtgggc gctcgcccct tgcccagtca     180
cactggtgtg tgtgcgactc ctacaaagtt aacagtttct ccaggtcaag gggtgggatc     240
caggcttggt gatgtgcaca atttcttttg tccacttgac acatctctgc gtcctgattc     300
tgctcaggga cggacccaag aacaaagcag ccatttaccg cctccggagg ggaggccagc     360
cctgtggcac atccagggcc ttggaacacc tagagacaga t                        401
```

<210> 36
<211> 401
<212> DNA
<213> Homo sapiens

<400> 36

```
gggtttcccc caagcccctt tccccctttg cgcctcccac ttctcctaga ttgagagtca      60
gcttggttct ttcctttaca tccattagtg agggtcaggc tcttttgtta tgtttttttt     120
tcttttgtat aacttaatta tttcagggtt cggggtgggc gctcgcccct tgcccagtca     180
cactggtgtg tgtgcgactc ttacaaagtt aacagtttct ccaggtcaag gggtgggatc     240
caggcttggt gatgtgcaca atttcttttg tccacttgac acatctctgc gtcctgattc     300
tgctcaggga cggacccaag aacaaagcag ccatttaccg cctccggagg ggaggccagc     360
cctgtggcac atccagggcc ttggaacacc tagagacaga t                        401
```

<210> 37
<211> 1068
<212> PRT
<213> Homo sapiens

<400> 37

```
Met Leu Val Gly Ser Gln Ser Phe Ser Pro Gly Gly Pro Asn Gly Ile
1               5                   10                  15
Ile Arg Ser Gln Ser Phe Ala Gly Phe Ser Gly Leu Gln Glu Arg Arg
            20                  25                  30
Ser Arg Cys Asn Ser Phe Ile Glu Asn Ser Ser Ala Leu Lys Lys Pro
        35                  40                  45
Gln Ala Lys Leu Lys Lys Met His Asn Leu Gly His Lys Asn Asn Asn
    50                  55                  60
Pro Pro Lys Glu Pro Gln Pro Lys Arg Val Glu Glu Val Tyr Arg Ala
65                  70                  75                  80
Leu Lys Asn Gly Leu Asp Glu Tyr Leu Glu Val His Gln Thr Glu Leu
                85                  90                  95
Asp Lys Leu Thr Ala Gln Leu Lys Asp Met Lys Arg Asn Ser Arg Leu
            100                 105                 110
Gly Val Leu Tyr Asp Leu Asp Lys Gln Ile Lys Thr Ile Glu Arg Tyr
        115                 120                 125
Met Arg Arg Leu Glu Phe His Ile Ser Lys Val Asp Glu Leu Tyr Glu
    130                 135                 140
Ala Tyr Cys Ile Gln Arg Arg Leu Gln Asp Gly Ala Ser Lys Met Lys
145                 150                 155                 160
Gln Ala Phe Ala Thr Ser Pro Ala Ser Lys Ala Ala Arg Glu Ser Leu
                165                 170                 175
```

```
Thr Glu Ile Asn Arg Ser Phe Lys Glu Tyr Thr Glu Asn Met Cys Thr
        180                 185                 190
Ile Glu Val Glu Leu Glu Asn Leu Leu Gly Glu Phe Ser Ile Lys Met
        195                 200                 205
Lys Gly Leu Ala Gly Phe Ala Arg Leu Cys Pro Gly Asp Gln Tyr Glu
    210                 215                 220
Ile Phe Met Lys Tyr Gly Arg Gln Arg Trp Lys Leu Lys Gly Lys Ile
225                 230                 235                 240
Glu Val Asn Gly Lys Gln Ser Trp Asp Gly Glu Glu Thr Val Phe Leu
            245                 250                 255
Pro Leu Ile Val Gly Phe Ile Ser Ile Lys Val Thr Glu Leu Lys Gly
            260                 265                 270
Leu Ala Thr His Ile Leu Val Gly Ser Val Thr Cys Glu Thr Lys Glu
        275                 280                 285
Leu Phe Ala Ala Arg Pro Gln Val Val Ala Val Asp Ile Asn Asp Leu
        290                 295                 300
Gly Thr Ile Lys Leu Asn Leu Glu Ile Thr Trp Tyr Pro Phe Asp Met
305                 310                 315                 320
Glu Asp Met Thr Ala Ser Ser Gly Ala Gly Asn Lys Ala Ala Ala Leu
            325                 330                 335
Gln Arg Arg Met Ser Met Tyr Ser Gln Gly Thr Pro Glu Thr Pro Thr
            340                 345                 350
Phe Lys Asp His Ser Phe Phe Arg Trp Leu His Pro Ser Pro Asp Lys
        355                 360                 365
Pro Arg Arg Leu Ser Val Leu Ser Ala Leu Gln Asp Thr Phe Phe Ala
    370                 375                 380
Lys Leu His Arg Ser Arg Ser Phe Ser Asp Leu Pro Ser Leu Arg Pro
385                 390             395                 400
Ser Pro Lys Ala Val Leu Glu Leu Tyr Ser Asn Leu Pro Asp Asp Ile
            405                 410                 415
Phe Glu Asn Gly Lys Ala Ala Glu Glu Lys Met Pro Leu Ser Leu Ser
            420                 425                 430
Phe Ser Asp Leu Pro Asn Gly Asp Cys Ala Leu Thr Ser His Ser Thr
        435                 440                 445
Gly Ser Pro Ser Asn Ser Thr Asn Pro Glu Ile Thr Ile Thr Pro Ala
    450                 455                 460
Glu Phe Asn Leu Ser Ser Leu Ala Ser Gln Asn Glu Gly Met Asp Asp
465                 470                 475                 480
Thr Ser Ser Ala Ser Ser Arg Asn Ser Leu Gly Glu Gly Gln Glu Pro
            485                 490                 495
Lys Ser His Leu Lys Glu Glu Asp Pro Glu Glu Pro Arg Lys Pro Ala
        500                 505                 510
Ser Ala Pro Ser Glu Ala Cys Arg Arg Gln Ser Ser Gly Ala Gly Ala
    515                 520                 525
Glu His Leu Phe Leu Glu Asn Asp Val Ala Glu Ala Leu Leu Gln Glu
    530                 535                 540
Ser Glu Glu Ala Ser Glu Leu Lys Pro Val Glu Leu Asp Thr Ser Glu
545                 550                 555                 560
Gly Asn Ile Thr Lys Gln Leu Val Lys Arg Leu Thr Ser Ala Glu Val
            565                 570                 575
Pro Met Ala Thr Asp Arg Leu Leu Ser Glu Gly Ser Val Gly Gly Glu
        580                 585                 590
Ser Glu Gly Cys Arg Ser Phe Leu Asp Gly Ser Leu Glu Asp Ala Phe
        595                 600                 605
Asn Gly Leu Leu Leu Ala Leu Glu Pro His Lys Glu Gln Tyr Lys Glu
    610                 615                 620
Phe Gln Asp Leu Asn Gln Glu Val Met Asn Leu Asp Asp Ile Leu Lys
```

```
625                     630                     635                     640
Cys Lys Pro Ala Val Ser Arg Ser Arg Ser Ser Ser Leu Ser Leu Thr
                645                     650                     655
Val Glu Ser Ala Leu Glu Ser Phe Asp Phe Leu Asn Thr Ser Asp Phe
                660                     665                     670
Asp Glu Glu Glu Asp Gly Asp Glu Val Cys Asn Val Gly Gly Gly Ala
            675                     680                     685
Asp Ser Val Phe Ser Asp Thr Glu Thr Glu Lys His Ser Tyr Arg Ser
        690                     695                     700
Val His Pro Glu Ala Arg Gly His Leu Ser Glu Ala Leu Thr Glu Asp
705                     710                     715                     720
Thr Gly Val Gly Thr Ser Val Ala Gly Ser Pro Leu Pro Leu Thr Thr
                725                     730                     735
Gly Asn Glu Ser Leu Asp Ile Thr Ile Val Arg His Leu Gln Tyr Cys
                740                     745                     750
Thr Gln Leu Val Gln Gln Ile Val Phe Ser Ser Lys Thr Pro Phe Val
                755                     760                     765
Ala Arg Ser Leu Leu Glu Lys Leu Ser Arg Gln Ile Gln Val Met Glu
        770                     775                     780
Lys Leu Ala Ala Val Ser Asp Glu Asn Ile Gly Asn Ile Ser Ser Val
785                     790                     795                     800
Val Glu Ala Ile Pro Glu Phe His Lys Lys Leu Ser Leu Leu Ser Phe
                805                     810                     815
Trp Thr Lys Cys Cys Ser Pro Val Gly Val Tyr His Ser Pro Ala Asp
                820                     825                     830
Arg Val Met Lys Gln Leu Glu Ala Ser Phe Ala Arg Thr Val Asn Lys
            835                     840                     845
Glu Tyr Pro Gly Leu Ala Asp Pro Val Phe Arg Thr Leu Val Ser Gln
        850                     855                     860
Ile Leu Asp Gln Ala Glu Pro Leu Leu Ser Ser Ser Leu Ser Ser Glu
865                     870                     875                     880
Val Val Thr Val Phe Gln Tyr Tyr Ser Tyr Phe Thr Ser His Gly Val
                885                     890                     895
Ser Asp Leu Glu Ser Tyr Leu Ser Gln Leu Ala Arg Gln Val Ser Met
            900                     905                     910
Val Gln Thr Leu Gln Ser Leu Arg Asp Glu Lys Leu Leu Gln Thr Met
        915                     920                     925
Ser Asp Leu Ala Pro Ser Asn Leu Leu Ala Gln Gln Glu Val Leu Arg
        930                     935                     940
Thr Leu Ala Leu Leu Leu Thr Arg Glu Asp Asn Glu Val Ser Glu Ala
945                     950                     955                     960
Val Thr Leu Tyr Leu Ala Ala Ala Ser Lys Asn Gln His Phe Arg Glu
                965                     970                     975
Lys Ala Leu Leu Tyr Tyr Cys Glu Ala Leu Thr Lys Thr Asn Leu Gln
            980                     985                     990
Leu Gln Lys Ala Ala Cys Leu Ala Leu Lys Ile Leu Glu Ala Thr Glu
        995                     1000                    1005
Ser Ile Lys Met Leu Val Thr Leu Cys Gln Ser Asp Thr Glu Glu Ile
    1010                    1015                    1020
Arg Asn Val Ala Ser Glu Thr Leu Leu Ser Leu Gly Glu Asp Gly Arg
1025                    1030                    1035                    1040
Leu Ala Tyr Glu Gln Leu Asp Lys Phe Pro Arg Asp Cys Val Lys Val
                1045                    1050                    1055
Gly Gly Arg His Gly Thr Glu Val Ala Thr Ala Phe
            1060                    1065
```

<210> 38

<211> 591
<212> PRT
<213> Homo sapiens

<400> 38
Met Leu Val Gly Ser Gln Ser Phe Ser Pro Gly Gly Pro Asn Gly Ile
1   5   10   15
Ile Arg Ser Gln Ser Phe Ala Gly Phe Ser Gly Leu Gln Glu Arg Arg
            20   25   30
Ser Arg Cys Asn Ser Phe Ile Glu Asn Ser Ser Ala Leu Lys Lys Pro
            35   40   45
Gln Ala Lys Leu Lys Lys Met His Asn Leu Gly His Lys Asn Asn Asn
        50   55   60
Pro Pro Lys Glu Pro Gln Pro Lys Arg Val Glu Glu Val Tyr Arg Ala
65   70   75   80
Leu Lys Asn Gly Leu Asp Glu Tyr Leu Glu Val His Gln Thr Glu Leu
                85   90   95
Asp Lys Leu Thr Ala Gln Leu Lys Asp Met Lys Arg Asn Ser Arg Leu
            100   105   110
Gly Val Leu Tyr Asp Leu Asp Lys Gln Ile Lys Thr Ile Glu Arg Tyr
        115   120   125
Met Arg Arg Leu Glu Phe His Ile Ser Lys Val Asp Glu Leu Tyr Glu
        130   135   140
Ala Tyr Cys Ile Gln Arg Arg Leu Gln Asp Gly Ala Ser Lys Met Lys
145   150   155   160
Gln Ala Phe Ala Thr Ser Pro Ala Ser Lys Ala Ala Arg Glu Ser Leu
                165   170   175
Thr Glu Ile Asn Arg Ser Phe Lys Glu Tyr Thr Glu Asn Met Cys Thr
                180   185   190
Ile Glu Val Glu Leu Glu Asn Leu Leu Gly Glu Phe Ser Ile Lys Met
        195   200   205
Lys Gly Leu Ala Gly Phe Ala Arg Leu Cys Pro Gly Asp Gln Tyr Glu
    210   215   220
Ile Phe Met Lys Tyr Gly Arg Gln Arg Trp Lys Leu Lys Gly Lys Ile
225   230   235   240
Glu Val Asn Gly Lys Gln Ser Trp Asp Gly Glu Glu Thr Val Phe Leu
                245   250   255
Pro Leu Ile Val Gly Phe Ile Ser Ile Lys Val Thr Glu Leu Lys Gly
                260   265   270
Leu Ala Thr His Ile Leu Val Gly Ser Val Thr Cys Glu Thr Lys Glu
        275   280   285
Leu Phe Ala Ala Arg Pro Gln Val Val Ala Val Asp Ile Asn Asp Leu
        290   295   300
Gly Thr Ile Lys Leu Asn Leu Glu Ile Thr Trp Tyr Pro Phe Asp Met
305   310   315   320
Glu Asp Met Thr Ala Ser Ser Gly Ala Gly Asn Lys Ala Ala Ala Leu
            325   330   335
Gln Arg Arg Met Ser Met Tyr Ser Gln Gly Thr Pro Glu Thr Pro Thr
            340   345   350
Phe Lys Asp His Ser Phe Phe Ser Asn Leu Pro Asp Asp Ile Phe Glu
        355   360   365
Asn Gly Lys Ala Ala Glu Glu Lys Met Pro Leu Ser Leu Ser Phe Ser
        370   375   380
Asp Leu Pro Asn Gly Asp Cys Ala Leu Thr Ser His Ser Thr Gly Ser
385   390   395   400
Pro Ser Asn Ser Thr Asn Pro Glu Ile Thr Ile Thr Pro Ala Glu Phe
                405   410   415

```
Asn Leu Ser Ser Leu Ala Ser Gln Asn Glu Gly Met Asp Asp Thr Ser
        420                 425                 430
Ser Ala Ser Ser Arg Asn Ser Leu Gly Glu Gly Gln Glu Pro Lys Ser
        435                 440                 445
His Leu Lys Glu Glu Asp Pro Glu Glu Pro Arg Lys Pro Ala Ser Ala
    450                 455                 460
Pro Ser Glu Ala Cys Arg Arg Gln Ser Ser Gly Ala Gly Ala Glu His
465                 470                 475                 480
Leu Phe Leu Glu Asn Asp Val Ala Glu Ala Leu Leu Gln Glu Ser Glu
                485                 490                 495
Glu Ala Ser Glu Leu Lys Pro Val Glu Leu Asp Thr Ser Glu Gly Asn
        500                 505                 510
Ile Thr Lys Gln Leu Val Lys Arg Leu Thr Ser Ala Glu Val Pro Met
        515                 520                 525
Ala Thr Asp Arg Leu Leu Ser Glu Gly Ser Val Gly Gly Glu Ser Glu
        530                 535                 540
Gly Cys Arg Ser Phe Leu Asp Gly Ser Leu Glu Asp Ala Phe Asn Gly
545                 550                 555                 560
Leu Leu Leu Ala Leu Glu Pro His Lys Glu Gln Tyr Lys Glu Phe Gln
                565                 570                 575
Asp Leu Asn Gln Glu Val Met Asn Leu Asp Asp Ile Leu Lys Lys
                580                 585                 590
```

<210> 39
<211> 1048
<212> PRT
<213> Homo sapiens

<400> 39

```
Met Leu Val Gly Ser Gln Ser Phe Ser Pro Gly Gly Pro Asn Gly Ile
 1               5                   10                  15
Ile Arg Ser Gln Ser Phe Ala Gly Phe Ser Gly Leu Gln Glu Arg Arg
                20                  25                  30
Ser Arg Cys Asn Ser Phe Ile Glu Asn Ser Ser Ala Leu Lys Lys Pro
        35                  40                  45
Gln Ala Lys Leu Lys Lys Met His Asn Leu Gly His Lys Asn Asn Asn
    50                  55                  60
Pro Pro Lys Glu Pro Gln Pro Lys Arg Val Glu Glu Val Tyr Arg Ala
65                  70                  75                  80
Leu Lys Asn Gly Leu Asp Glu Tyr Leu Glu Val His Gln Thr Glu Leu
                85                  90                  95
Asp Lys Leu Thr Ala Gln Leu Lys Asp Met Lys Arg Asn Ser Arg Leu
        100                 105                 110
Gly Val Leu Tyr Asp Leu Asp Lys Gln Ile Lys Thr Ile Glu Arg Tyr
        115                 120                 125
Met Arg Arg Leu Glu Phe His Ile Ser Lys Val Asp Glu Leu Tyr Glu
    130                 135                 140
Ala Tyr Cys Ile Gln Arg Arg Leu Gln Asp Gly Ala Ser Lys Met Lys
145                 150                 155                 160
Gln Ala Phe Ala Thr Ser Pro Ala Ser Lys Ala Ala Arg Glu Ser Leu
                165                 170                 175
Thr Glu Ile Asn Arg Ser Phe Lys Glu Tyr Thr Glu Asn Met Cys Thr
                180                 185                 190
Ile Glu Val Glu Leu Glu Asn Leu Leu Gly Glu Phe Ser Ile Lys Met
        195                 200                 205
Lys Gly Leu Ala Gly Phe Ala Arg Leu Cys Pro Gly Asp Gln Tyr Glu
    210                 215                 220
```

```
Ile Phe Met Lys Tyr Gly Arg Gln Arg Trp Lys Leu Lys Gly Lys Ile
225                 230                 235                 240
Glu Val Asn Gly Lys Gln Ser Trp Asp Gly Glu Glu Thr Val Phe Leu
                245                 250           ,         255
Pro Leu Ile Val Gly Phe Ile Ser Ile Lys Val Thr Glu Leu Lys Gly
                260                 265                 270
Leu Ala Thr His Ile Leu Val Gly Ser Val Thr Cys Glu Thr Lys Glu
                275                 280                 285
Leu Phe Ala Ala Arg Pro Gln Val Val Ala Val Asp Ile Asn Asp Leu
                290                 295                 300
Gly Thr Ile Lys Leu Asn Leu Glu Ile Thr Trp Tyr Pro Phe Asp Met
305                 310                 315                 320
Glu Asp Met Thr Ala Ser Ser Gly Ala Gly Asn Lys Ala Ala Ala Leu
                325                 330                 335
Gln Arg Arg Met Ser Met Tyr Ser Gln Gly Thr Pro Glu Thr Pro Thr
                340                 345                 350
Phe Lys Asp His Ser Phe Phe Arg Trp Leu His Pro Ser Pro Asp Lys
                355                 360                 365
Pro Arg Arg Leu Ser Val Leu Ser Ala Leu Gln Asp Thr Phe Phe Ala
                370                 375                 380
Lys Leu His Arg Ser Arg Ser Phe Ser Asp Leu Pro Ser Leu Arg Pro
385                 390                 395                 400
Ser Pro Lys Ala Val Leu Glu Leu Tyr Ser Asn Leu Pro Asp Asp Ile
                405                 410                 415
Phe Glu Asn Gly Lys Ala Ala Glu Glu Lys Met Pro Leu Ser Leu Ser
                420                 425                 430
Phe Ser Asp Leu Pro Asn Gly Asp Cys Ala Leu Thr Ser His Ser Thr
                435                 440                 445
Gly Ser Pro Ser Asn Ser Thr Asn Pro Glu Ile Thr Ile Thr Pro Ala
450                 455                 460
Glu Phe Asn Leu Ser Ser Leu Ala Ser Gln Asn Glu Gly Met Asp Asp
465                 470                 475                 480
Thr Ser Ser Ala Ser Ser Arg Asn Ser Leu Gly Glu Gly Gln Glu Pro
                485                 490                 495
Lys Ser His Leu Lys Glu Glu Asp Pro Glu Glu Pro Arg Lys Pro Ala
                500                 505                 510
Ser Ala Pro Ser Glu Ala Cys Arg Arg Gln Ser Ser Gly Ala Gly Ala
                515                 520                 525
Glu His Leu Phe Leu Glu Asn Asp Val Ala Glu Ala Leu Leu Gln Glu
                530                 535                 540
Ser Glu Glu Ala Ser Glu Leu Lys Pro Val Glu Leu Asp Thr Ser Glu
545                 550                 555                 560
Gly Asn Ile Thr Lys Gln Leu Val Lys Arg Leu Thr Ser Ala Glu Val
                565                 570                 575
Pro Met Ala Thr Asp Arg Leu Leu Ser Glu Gly Ser Val Gly Gly Glu
                580                 585                 590
Ser Glu Gly Cys Arg Ser Phe Leu Asp Gly Ser Leu Glu Asp Ala Phe
                595                 600                 605
Asn Gly Leu Leu Leu Ala Leu Glu Pro His Lys Glu Gln Tyr Lys Glu
                610                 615                 620
Phe Gln Asp Leu Asn Gln Glu Val Met Asn Leu Asp Asp Ile Leu Lys
625                 630                 635                 640
Cys Lys Pro Ala Val Ser Arg Ser Arg Ser Ser Ser Leu Ser Leu Thr
                645                 650                 655
Val Glu Ser Ala Leu Glu Ser Phe Asp Phe Leu Asn Thr Ser Asp Phe
                660                 665                 670
Asp Glu Glu Glu Asp Gly Asp Glu Val Cys Asn Val Gly Gly Gly Ala
```

```
              675                    680                    685
Asp Ser Val Phe Ser Asp Thr Glu Thr Glu Lys His Ser Tyr Arg Ser
     690                    695                    700
Val His Pro Glu Ala Arg Gly His Leu Ser Glu Ala Leu Thr Glu Asp
705                    710                    715                    720
Thr Gly Val Gly Thr Ser Val Ala Gly Ser Pro Leu Pro Leu Thr Thr
               725                    730                    735
Gly Asn Glu Ser Leu Asp Ile Thr Ile Val Arg His Leu Gln Tyr Cys
               740                    745                    750
Thr Gln Leu Val Gln Gln Ile Val Phe Ser Ser Lys Thr Pro Phe Val
               755                    760                    765
Ala Arg Ser Leu Leu Glu Lys Leu Ser Arg Gln Ile Gln Val Met Glu
          770                    775                    780
Lys Leu Ala Ala Val Ser Asp Glu Asn Ile Gly Asn Ile Ser Ser Val
785                    790                    795                    800
Val Glu Ala Ile Pro Glu Phe His Lys Lys Leu Ser Leu Leu Ser Phe
               805                    810                    815
Trp Thr Lys Cys Cys Ser Pro Val Gly Val Tyr His Ser Pro Ala Asp
          820                    825                    830
Arg Val Met Lys Gln Leu Glu Ala Ser Phe Ala Arg Thr Val Asn Lys
          835                    840                    845
Glu Tyr Pro Gly Leu Ala Asp Pro Val Phe Arg Thr Leu Val Ser Gln
     850                    855                    860
Ile Leu Asp Gln Ala Glu Pro Leu Leu Ser Ser Ser Leu Ser Ser Glu
865                    870                    875                    880
Val Val Thr Val Phe Gln Tyr Tyr Ser Tyr Phe Thr Ser His Gly Val
               885                    890                    895
Ser Asp Leu Glu Ser Tyr Leu Ser Gln Leu Ala Arg Gln Val Ser Met
          900                    905                    910
Val Gln Thr Leu Gln Ser Leu Arg Asp Glu Lys Leu Leu Gln Thr Met
     915                    920                    925
Ser Asp Leu Ala Pro Ser Asn Leu Leu Ala Gln Gln Glu Val Leu Arg
     930                    935                    940
Thr Leu Ala Leu Leu Leu Thr Arg Glu Asp Asn Glu Val Ser Glu Ala
945                    950                    955                    960
Val Thr Leu Tyr Leu Ala Ala Ala Ser Lys Asn Gln His Phe Arg Glu
               965                    970                    975
Lys Ala Leu Leu Tyr Tyr Cys Glu Ala Leu Thr Lys Thr Asn Leu Gln
               980                    985                    990
Leu Gln Lys Ala Ala Cys Leu Ala Leu Lys Ile Leu Glu Ala Thr Glu
          995                    1000                   1005
Ser Ile Lys Met Leu Val Thr Leu Cys Gln Ser Asp Thr Glu Glu Ile
     1010                   1015                   1020
Arg Asn Val Ala Ser Glu Thr Leu Leu Ser Leu Gly Glu Asp Gly Arg
1025                   1030                   1035                   1040
Leu Ala Tyr Glu Gln Leu Asp Lys
               1045
```

```
<210> 40
<211> 479
<212> PRT
<213> Homo sapiens

<400> 40
Met Leu Val Gly Ser Gln Ser Phe Ser Pro Gly Gly Pro Asn Gly Ile
 1               5                    10                   15
Ile Arg Ser Gln Ser Phe Ala Gly Phe Ser Gly Leu Gln Glu Arg Arg
```

```
                    20                      25                      30
Ser Arg Cys Asn Ser Phe Ile Glu Asn Ser Ser Ala Leu Lys Lys Pro
             35                      40                      45
Gln Ala Lys Leu Lys Lys Met His Asn Leu Gly His Lys Asn Asn Asn
    50                      55                      60.
Pro Pro Lys Glu Pro Gln Pro Lys Arg Val Glu Glu Val Tyr Arg Ala
65                      70                      75                      80
Leu Lys Asn Gly Leu Asp Glu Tyr Leu Glu Val His Gln Thr Glu Leu
                85                      90                      95
Asp Lys Leu Thr Ala Gln Leu Lys Asp Met Lys Arg Asn Ser Arg Leu
                100                     105                     110
Gly Val Leu Tyr Asp Leu Asp Lys Gln Ile Lys Thr Ile Glu Arg Tyr
                115                     120                     125
Met Arg Arg Leu Glu Phe His Ile Ser Lys Val Asp Glu Leu Tyr Glu
                130                     135                     140
Ala Tyr Cys Ile Gln Arg Arg Leu Gln Asp Gly Ala Ser Lys Met Lys
145                     150                     155                     160
Gln Ala Phe Ala Thr Ser Pro Ala Ser Lys Ala Ala Arg Glu Ser Leu
                165                     170                     175
Thr Glu Ile Asn Arg Ser Phe Lys Glu Tyr Thr Glu Asn Met Cys Thr
                180                     185                     190
Ile Glu Val Glu Leu Glu Asn Leu Leu Gly Glu Phe Ser Ile Lys Met
                195                     200                     205
Lys Gly Leu Ala Gly Phe Ala Arg Leu Cys Pro Gly Asp Gln Tyr Glu
    210                     215                     220
Ile Phe Met Lys Tyr Gly Arg Gln Arg Trp Lys Leu Lys Gly Lys Ile
225                     230                     235                     240
Glu Val Asn Gly Lys Gln Ser Trp Asp Gly Glu Glu Thr Val Phe Leu
                245                     250                     255
Pro Leu Ile Val Gly Phe Ile Ser Ile Lys Val Thr Glu Leu Lys Gly
                260                     265                     270
Leu Ala Thr His Ile Leu Val Gly Ser Val Thr Cys Glu Thr Lys Glu
                275                     280                     285
Leu Phe Ala Ala Arg Pro Gln Val Val Ala Val Asp Ile Asn Asp Leu
                290                     295                     300
Gly Thr Ile Lys Leu Asn Leu Glu Ile Thr Trp Tyr Pro Phe Asp Met
305                     310                     315                     320
Glu Asp Met Thr Ala Ser Ser Gly Ala Gly Asn Lys Ala Ala Ala Leu
                325                     330                     335
Gln Arg Arg Met Ser Met Tyr Ser Gln Gly Thr Pro Glu Thr Pro Thr
                340                     345                     350
Phe Lys Asp His Ser Phe Phe Ser Asn Leu Pro Asp Asp Ile Phe Glu
                355                     360                     365
Asn Gly Lys Ala Ala Glu Glu Lys Met Pro Leu Ser Leu Ser Phe Ser
    370                     375                     380
Asp Leu Pro Asn Gly Asp Cys Ala Leu Thr Ser His Ser Thr Gly Ser
385                     390                     395                     400
Pro Ser Asn Ser Thr Asn Pro Glu Ile Thr Ile Thr Pro Ala Glu Phe
                405                     410                     415
Asn Leu Ser Ser Leu Ala Ser Gln Asn Glu Gly Met Asp Asp Thr Ser
                420                     425                     430
Ser Ala Ser Ser Arg Asn Ser Leu Gly Glu Gly Gln Glu Pro Lys Ser
                435                     440                     445
His Leu Lys Glu Glu Asp Pro Glu Glu Pro Arg Lys Pro Ala Ser Ala
    450                     455                     460
Pro Ser Glu Ala Cys Arg Arg Gln Ser Ser Gly Ala Gly Ala Glu
465                     470                     475
```